PROF. DR. CHASE BEISEL

Group Leader

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Education

2009	PhD, Chemical Engineering, California Institute of Technology, Pasadena, CA, USA
2004	BSc, Chemical Engineering, Iowa State University, Ames, IA, USA

Positions

2018 - present	Group Leader, Helmholtz Institute for RNA-based Infection Research (HIRI), Würzburg, Germany
2018 - present	Professor (W2), Faculty of Medicine, University of Würzburg, Germany
2017 - 2018	Associate Professor, Department of Chemical and Biomolecular Engineering, North Carolina
2015 - present 2011 - 2017 2009 - 2011	State University (NCSU), Raleigh, NC, USA Co-founder and scientific advisory board member, Locus Biosciences, Morrisville, NC, USA Assistant Professor, Dept. of Chemical and Biomolecular Engineering, NCSU, Raleigh, NC, USA Postdoctoral fellow, National Institutes of Health, Bethesda, MD, USA

Committee Work

2019	Organizer, International Conference on CRISPR Technologies, Würzburg, Germany
2017	Organizer, International Conference on CRISPR Technologies, Raleigh, NC, USA
2016 - 2017	Instructor, Cold Spring Harbor Laboratory Synthetic Biology Summer School, NY, USA
2016	Topical A Conference Chair, AIChE Annual Conference, San Francisco, CA, USA

Awards & Honors

AlChE Program Development Service Award (2018), D.I.C. Wang Young Investigator Award (2018), Camille Dreyfus Teacher-Scholar Award (2017), Bay Area Lyme Foundation Emerging Leader Award (2016), Sigma Xi Faculty Research Award (2016), NCSU Faculty Scholar (2015 - 2016), NSF CAREER Award (2015), Gordon & Betty Moore Foundation Fellow, Life Sciences Research Foundation (2010 - 2011)

Current Boards

Scientific Advisory Board at Locus Biosciences

Selected Publications

Dugar G, Leenay RT, Eisenbart SK, Bischler T, Aul BU, **Beisel CL**, Sharma CM (2018) *CRISPR RNA-dependent binding and cleavage of endogenous RNAs by the Campylobacter jejuni Cas9* **Molecular Cell** 69(5): p893–905.e7

Marshall R, Maxwell CS, Collins SP, Jacobsen T, Luo ML, Begemann MB, Gray BN, January E, Singer A, He Y, **Beisel CL**, Noireaux V (2018) Rapid and Scalable Characterization of CRISPR Technologies Using an E. coli Cell-Free Transcription-Translation System **Molecular Cell** 69(1):146-157

Leenay RT, Maksimchuk KR, Slotkowski RA, Agrawal RN, Gomaa AA, Briner AE, Barrangou R, **Beisel CL** (2016) *Identifying and Visualizing Functional PAM Diversity across CRISPR-Cas Systems* **Molecular Cell** 62(1):137-47

Luo ML, Mullis AS, Leenay RT, **Beisel CL** (2015) *Repurposing endogenous type I CRISPR-Cas systems for programmable gene repression* **Nucleic Acids Research** 43(1):674-81

Gomaa AA, Klumpe HE, Luo ML, Selle K, Barrangou R, **Beisel CL** (2014) *Programmable removal of bacterial strains by use of genome-targeting CRISPR-Cas systems* **mBio** 5(1):e00928-13